

## AMENDMENTS TO THE SPECIFICATION

**On page 6, please replace the paragraph beginning on line 10 and ending on line 15 with the following amended paragraph:**

Figure 4 shows Northern and RACE Analysis of MCFD2. Figure 4A, Northern blot of polyA mRNA (2 mg/lane) from multiple human tissues was hybridized with a probe spanning the entire coding region of MCFD2. A 4.1-kb mRNA can be seen in all the tissues, although faint in brain and lung. Smaller transcripts in the range of 0.8~1.8-kb are also detected in some tissues. Figure 4B (SEQ ID NOS:29-31), the 5' untranslated region of the MCFD2 gene. Arrows indicate transcriptional start sites as determined by 5' RACE.

**On page 89, please replace the paragraph beginning on line 25 and ending on page 90, line 8 with the following amended paragraph:**

Total cellular RNA was prepared from EBV-transformed lymphoblasts and HeLa cells. The 5' end of the mRNA was determined by rapid amplification of cDNA ends (RACE) with a FirstChoice RLM-RACE kit (Ambion) using primers AGCAGGCCACACAGGAAG (SEQ ID NO: 27) and CTCTTGGTCGTGCACTGTGT (SEQ ID NO:3). Sequence of RTPCS products was determined by the University of Michigan DNA Sequencing Core. A PCR product containing the coding sequence of the MCFD2 cDNA was amplified from a human MOLT4 T-cell cDNA library as previously described (Levy et al., 2001) using primers GCTTGGTACCTGCAGTGATTTTGCAAATTCAG (SEQ ID NO: 28) and GGACTCGAGACCATGAGATCCCTGCTCAGA (SEQ ID NO 4). After gel purification, this DNA fragment was labeled by random priming and used to hybridize to a FirstChoice Human Northern Blot (Ambion) in Rapid-Hyb buffer (Amersham) according to the manufacturer's specifications. The same pair of primers was used to screen MCFD2 expression by PCR on cDNA obtained from a multiple tissue cDNA panel (Clontech).

Please insert the attached Sequence Listing as new pages --106-218--.

**IN THE CLAIMS**

Please renumber the Claims pages from pages "106-107" to --219-220--.

**IN THE ABSTRACT:**

Please renumber the Abstract page from page "108" to --221--.